

```
Fig. 1
                                                                                                    Tn7
                                                                  TGTTTAAACACAAT--Kanr--ATTGTGTTTAAACA
-- N_1N_2N_3N_4N_5N_6N_7N_8N_9N_{10}---
                                                                  ACAAATTTGTGTTA--Kanr--TAACACAAATTTGT
-- N_1N_2N_3N_4N_5N_6N_7N_8N_9N_{10}---
                                                                                                  (SEQ ID NO.: 10)
 Step 1
(a) Insertion of transposon into target DNA
-- N<sub>1</sub>N<sub>2</sub>N<sub>3</sub>N<sub>4</sub>N<sub>5</sub>N<sub>6</sub> ACAAATTTGTGTTA--Kan<sup>r</sup>--TAACACAAATTTGT N<sub>2</sub>N<sub>3</sub>N<sub>4</sub>N<sub>5</sub>N<sub>6</sub>N<sub>7</sub>N<sub>8</sub>N<sub>9</sub>N<sub>10</sub>---
                                     (SEQ ID NO.: 10)
(b) Removal of transposon
                                                                         Pme I
              -- N_1N_2N_3N_4N_5N_6TGTTT
                                                                                  AAACAN_2N_3N_4N_5N_6N_7N_8N_9N_{10} - - -
                                                                                  TTTGTN_2N_3N_4N_5N_6N_7N_8N_9N_{10}---
              -- N<sub>1</sub>N<sub>2</sub>N<sub>3</sub>N<sub>4</sub>N<sub>5</sub>N<sub>6</sub>ACAAA
 Step 2
(a) Insertion of a first cassette into the cut site
                                       -- N<sub>1</sub>N<sub>2</sub>N<sub>3</sub>N<sub>4</sub>N<sub>5</sub>N<sub>6</sub>TGTTTGGATCC-----GTGCAG AATGCCAAACAN<sub>2</sub>N<sub>3</sub>N<sub>4</sub>N<sub>5</sub>N<sub>6</sub>N<sub>7</sub>N<sub>8</sub>N<sub>9</sub>N<sub>10</sub>---
  -- N<sub>1</sub>N<sub>2</sub>N<sub>3</sub>N<sub>4</sub>N<sub>5</sub>N<sub>6</sub>ACAAACCTAGG------CACGTC TTACGGTTTGTN<sub>2</sub>N<sub>3</sub>N<sub>4</sub>N<sub>5</sub>N<sub>6</sub>N<sub>7</sub>N<sub>8</sub>N<sub>9</sub>N<sub>10</sub>
                                                                      (SEQ ID NO.: 4)
                                                                          Bsg I, T4 DNA Polymerase
(b) Removal of the first cassette
                                                                          BamH I (Removal of 3'-overhang)
              -- N<sub>1</sub>N<sub>2</sub>N<sub>3</sub>N<sub>4</sub>N<sub>5</sub>N<sub>6</sub>TGTTTG
                                                                                      N_7N_8N_9N_{10}---
              -- N<sub>1</sub>N<sub>2</sub>N<sub>3</sub>N<sub>4</sub>N<sub>5</sub>N<sub>6</sub>ACAAACCTAG
                                                                                      N_7 N_8 N_9 N_{10} - - -
  Step 3
 (a) Insertion of a second cassette into the cut site
                                   (SEQ ID NO.: 5)
-- N<sub>1</sub>N<sub>2</sub>N<sub>3</sub>N<sub>4</sub>N<sub>5</sub>N<sub>6</sub>TGTTTGGATCCTT CTGCAC---GTGCAG GATCCGTAATTGACGTATGN<sub>7</sub>N<sub>8</sub>N<sub>9</sub>N<sub>10</sub>-
-- N<sub>1</sub>N<sub>2</sub>N<sub>3</sub>N<sub>4</sub>N<sub>5</sub>N<sub>6</sub>ACAAACCTAGGAA GACGTG---CACGTC CTAGGCATTAACTGCATACN<sub>7</sub>N<sub>8</sub>N<sub>9</sub>N<sub>10</sub>--
                                                                                       (SEQ ID NO.: 6)
                                                                          Bsg I, T4 DNA Polymerase
 (b) Removal of the second cassette
                                                                          (Removal of 3'-overhang)
                                                                                          ATGN7N8N9N10---
                             --N_1N_2N_3
                                                                                           TACN7N8N9N10---
                             --N_1N_2N_3
                                                   --- N_1N_2N_3ATGN_7N_8N_9N_{10}---
                                                                                                           (Substitution of N<sub>4</sub>N<sub>5</sub>N<sub>6</sub>
 Step 4:
                                                 --- N_1N_2N_3 TACN_7N_8N_9N_{10} ---
                                                                                                           sequence with ATG)
 Self-ligation of the cut sites
```

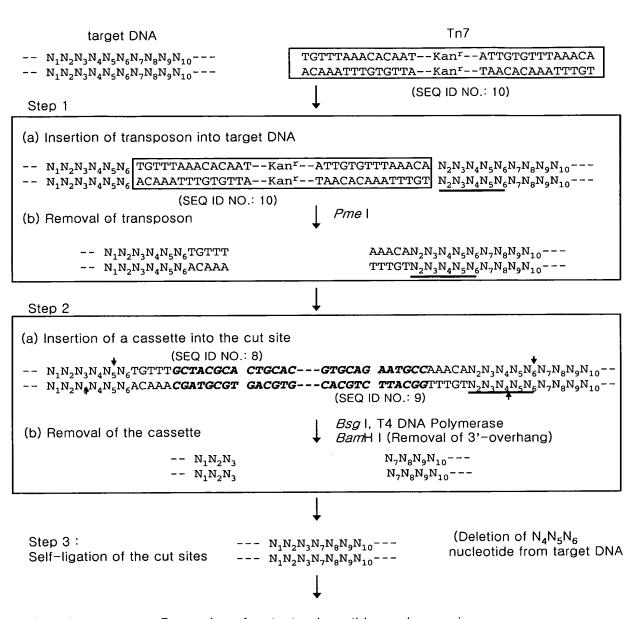
Step 5:

Expression of mutant polypeptides and screening

Fia. 2

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Tn7
                               target DNA
                                                                                                                            TGTTTAAACACAAT--Kanr--ATTGTGTTTAAACA
   -- N_1N_2N_3N_4N_5N_6N_7N_8N_9N_{10}---
                                                                                                                            ACAAATTTGTGTTA--Kan<sup>r</sup>--TAACACAAATTTGT
   -- N_1N_2N_3N_4N_5N_6N_7N_8N_9N_{10}---
                                                                                                                                                                   (SEQ ID NO.: 10)
    Step 1
  (a) Insertion of transposon into target DNA
  -- \text{ N}_1 \text{N}_2 \text{N}_3 \text{N}_4 \text{N}_5 \text{N}_6 \boxed{\text{TGTTTAAACACAAT} -\text{Kan}^r - \text{ATTGTGTTTAAACA}} \text{ N}_2 \text{N}_3 \text{N}_4 \text{N}_5 \text{N}_6 \text{N}_7 \text{N}_8 \text{N}_9 \text{N}_{10} ---
  -- \text{ N}_1 \text{ N}_2 \text{ N}_3 \text{ N}_4 \text{ N}_5 \text{ N}_6 \boxed{\text{ACAAATTTGTGTTA} - \text{Kan}^r - \text{TAACACAAATTTGT}} \text{ N}_2 \text{ N}_3 \text{ N}_4 \text{ N}_5 \text{ N}_6 \text{ N}_7 \text{ N}_8 \text{ N}_9 \text{ N}_{10} - --
                                                                         (SEQ ID NO.: 10)
                                                                                                                                        Pme I
  (b) Removal of transposon
                           -- N_1N_2N_3N_4N_5N_6TGTTT
                                                                                                                                                       AAACAN_2N_3N_4N_5N_6N_7N_8N_9N_{10} - - -
                                                                                                                                                       {\tt TTTGTN}_2{\tt N}_3{\tt N}_4{\tt N}_5{\tt N}_6{\tt N}_7{\tt N}_8{\tt N}_9{\tt N}_{10}{\tt ---}
                           -- N<sub>1</sub>N<sub>2</sub>N<sub>3</sub>N<sub>4</sub>N<sub>5</sub>N<sub>6</sub>ACAAA
    Step 2
  (a) Insertion of a first cassette into the cut site
                                                                        ♦ (SEQ ID NO.: 3)
      -- N<sub>1</sub>N<sub>2</sub>N<sub>3</sub>N<sub>4</sub>N<sub>5</sub>N<sub>6</sub>TGTTTGGATCC------GTGCAG AATGCCAAACAN<sub>2</sub>N<sub>3</sub>N<sub>4</sub>N<sub>5</sub>N<sub>6</sub>N<sub>7</sub>N<sub>8</sub>N<sub>9</sub>N<sub>10</sub>---
      -- N_1 N_2 N_3 N_4 N_5 N_6 A CAAA \textit{CCTAGG-------CACGTC} \quad \textit{TTACGG} \text{TTTGTN}_2 N_3 N_4 N_5 N_6 N_7 N_8 N_9 N_{10} - N_1 N_2 N_3 N_4 N_5 N_6 N_7 N_8 N_9 N_{10} - N_1 N_2 N_3 N_4 N_5 N_6 N_7 N_8 N_9 N_{10} - N_1 N_2 N_3 N_4 N_5 N_6 N_7 N_8 N_9 N_{10} - N_1 N_2 N_3 N_4 N_5 N_6 N_7 N_8 N_9 N_{10} - N_1 N_2 N_3 N_4 N_5 N_6 N_7 N_8 N_9 N_{10} - N_1 N_2 N_3 N_4 N_5 N_6 N_7 N_8 N_9 N_{10} - N_1 N_2 N_3 N_4 N_5 N_6 N_7 N_8 N_9 N_{10} - N_1 N_2 N_8 N_9 N_{10} N_9 N_{
                                                                                                                                  (SEQ ID NO.: 4)
                                                                                                                                   Bsg I, T4 DNA Polymerase
   (b) Removal of the first cassette
                                                                                                                                    BamH | (Removal of 3'-overhang)
                           -- N_1N_2N_3N_4N_5N_6TGTTTG
                                                                                                                                                              N_7 N_8 N_9 N_{10} - - -
                           -- N<sub>1</sub>N<sub>2</sub>N<sub>3</sub>N<sub>4</sub>N<sub>5</sub>N<sub>6</sub>ACAAACCTAG
                                                                                                                                                              N_7N_8N_9N_{10} ---
     Step 3
    (a) Insertion of a second cassette into the cut site
                                                                     (SEQ ID NO.: 7)
  - N<sub>1</sub>N<sub>2</sub>N<sub>3</sub>N<sub>4</sub>N<sub>5</sub>N<sub>6</sub>TGTTTGGATCCTTGCA CTGCAC--GTGCAG GATCCGTAATTGACGTNNNN<sub>7</sub>N<sub>8</sub>N<sub>9</sub>N<sub>10</sub>-
-- N<sub>1</sub>N<sub>2</sub>N<sub>3</sub>N<sub>4</sub>N<sub>5</sub>N<sub>6</sub>ACAAACCTAGGAACGT GACGTG--CACGTC CTAGGCATTAACTGCANNNN<sub>7</sub>N<sub>8</sub>N<sub>9</sub>N<sub>10</sub>-
                                                                                                                                                               (SEQ ID NO.: 6)
                                                                                                                                         Bsg I, T4 DNA Polymerase
   (b) Removal of the second cassette
                                                                                                                                         (Removal of 3'-overhang)
                                                                                                                                                                       NNN7N8N9N10---
                                      -- N_1 N_2 N_3 N_4 N_5 N_6
                                      --N_1N_2N_3N_4N_5N_6
                                                                                                                                                                       NNNN<sub>7</sub>N<sub>8</sub>N<sub>9</sub>N<sub>10</sub>---
                                                                                   (Insertion of random
   Self-ligation of the cut sites _{--} _{N_1N_2N_3N_4N_5N_6} _{N_0N_7N_8N_9N_{10}} _{N_1N_2N_3N_4N_5N_6}
                                                                                                                                                                                                        NNN sequence)
      Step 5:
                                                                     Expression of mutant polypeptides and screening
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Fig. 3



Step 4: Expression of mutant polypeptides and screening

GGTGA (N) 8^ CCACT (N) 7^	GAAGA (N) ₈ ^ CTTCT (N) ₇ ^	GAGTC (N) ₅ ^ CTCAG (N) ₅ ^	GAGTC (N) 4 ^ CTCAG (N) 5 ^	$GCTCTTC (N)_{1}^{\wedge}$ $CGAGAAG (N)_{4}^{\wedge}$	GCATC (N) 5^ CGTAG (N) 9^	GGATG (N) ₉ ^ CCTAC (N) ₁₃ ^	GACGC (N) ₅ ^ CTGCG (N) ₁₀ ^	
IqdH	Mboll	MlyI	PleI	SapI	SfaN1	FokI	HgaI	
GTCTC (N) 1 CAGAG (N) 5 CAGAG (N) 5 CAGAG (N)	CGTCTC (N) ₁ ^ GCAGAG (N) ₅ ^	$GGGAC\left(\mathrm{N}\right)_{10}^{\diamond}$ $CCCTG\left(\mathrm{N}\right)_{14}^{\diamond}$	ACCTGC (N) 4 ^ TGGACG (N) 8 ^	GCAATG $(N)_2$ CGTTAC $^{\circ}(N)_2$	GCAGTG $(N)_2$ CGTCAC $^{\prime}$ $(N)_2$	$\mathtt{CTCTTC}(\mathtt{N})_{1}^{\wedge}$ $\mathtt{GAGAAG}(\mathtt{N})_{4}^{\wedge}$	GGCGGA (N) 11 CCGCCT (N) 9 C	
BsmAI	BsmB1	BsmF1	BspM1	BsrDI	BtsI	EarI	EciI	
GGATC (N) 4 CCTAG (N) 5 C	GAAGAC (N) 2^ CTTCTG (N) 6^	GCAGC (N) ₈ ^ CGTCG (N) ₁₂ ^	GTATCC (N) ₆ ^ CATAGG (N) ₅ ^	ACTGGG (N) ₅ ^ TGACCC (N) ₄ ^	CTGGAG $\left(\mathrm{N}\right)_{16}^{\circ}$ GACCTC $\left(\mathrm{N}\right)_{14}^{\circ}$	GGTCTC (N) ₁ ^ CCAGAG (N) ₅ ^	GAGGAG (N) ₁₀ ^ CTCCTC (N) ₈ ^	GTGCAG (N) $_{16}^{\circ}$ CACGTC (N) $_{14}^{\circ}$
AlwI	BbsI	BbvI	BciVI	BmrI	BpmI	BsaI	BseRI	BsgI